

FILE COPY

Naval Research Laboratory

Washington, DC 20375-5000



NRL Memorandum Report 6254

Subroutine Probdif

G. V. TRUNK

*Radar Analysis Branch
Radar Division*

September 12, 1988

DTIC
ELECTE
OCT 11 1988
S D
D

Approved for public release; distribution unlimited.

88 10 6 T10

AD-A199 541

| REPORT DOCUMENTATION PAGE | | | | Form Approved OMB No. 0704-0188 | |
|--|-------|--|---|---|--------------------------------------|
| 1a REPORT SECURITY CLASSIFICATION UNCLASSIFIED | | | 1b RESTRICTIVE MARKINGS | | |
| 2a SECURITY CLASSIFICATION AUTHORITY | | | 3 DISTRIBUTION AVAILABILITY OF REPORT Approved for public release; distribution unlimited. | | |
| 2b DECLASSIFICATION/DOWNGRADING SCHEDULE | | | 5 MONITORING ORGANIZATION REPORT NUMBER(S) | | |
| 4 PERFORMING ORGANIZATION REPORT NUMBER(S) NRL Memorandum Report 6254 | | | | | |
| 6a NAME OF PERFORMING ORGANIZATION Naval Research Laboratory | | 6b OFFICE SYMBOL (If applicable) Code 5310 | 7a NAME OF MONITORING ORGANIZATION | | |
| 6c ADDRESS (City, State, and ZIP Code) Washington, DC 20375-5000 | | | 7b ADDRESS (City, State, and ZIP Code) | | |
| 8a NAME OF FUNDING/SPONSORING ORGANIZATION Naval Sea Systems Command | | 8b OFFICE SYMBOL (If applicable) | 9 PROCUREMENT INSTRUMENT IDENTIFICATION NUMBER | | |
| 8c ADDRESS (City, State, and ZIP Code) Washington, DC 20362-5101 | | | 10 SOURCE OF FUNDING NUMBERS | | |
| | | | PROGRAM ELEMENT NO 64573N | PROJECT NO S0954 | TASK NO WORK UNIT ACCESSION NO |
| 11 TITLE (Include Security Classification) Subroutine Probdif | | | | | |
| 12 PERSONAL AUTHOR(S) Trunk, G.V. | | | | | |
| 13a TYPE OF REPORT Interim | | 13b TIME COVERED FROM 3/88 TO 5/88 | | 14 DATE OF REPORT (Year, Month, Day) 1988 September 12 | |
| 15 PAGE COUNT 19 | | | | | |
| 16 SUPPLEMENTARY NOTATION | | | | | |
| 17 COSATI CODES | | | 18 SUBJECT TERMS (Continue on reverse if necessary and identify by block number) | | |
| FIELD | GROUP | SUB-GROUP | Distributions | | |
| | | | Ordered samples | | |
| 19 ABSTRACT (Continue on reverse if necessary and identify by block number) | | | | | |
| <p>This reports describes a technique for determining the probability that one density is a shifted version of another density. We assume nothing about the density and employ a nonparametric procedure on the sample data.</p> | | | | | |
| 20 DISTRIBUTION AVAILABILITY OF ABSTRACT <input checked="" type="checkbox"/> UNCLASSIFIED/UNLIMITED <input type="checkbox"/> SAME AS RPT <input type="checkbox"/> DTIC USERS | | | 21 ABSTRACT SECURITY CLASSIFICATION UNCLASSIFIED | | |
| 22a NAME OF RESPONSIBLE INDIVIDUAL G.V. Trunk | | | 22b TELEPHONE (include Area Code) : 22c OFFICE SYMBOL (202) 767-2573 Code 5310 | | |

CONTENTS

| | |
|---|---|
| INTRODUCTION | 1 |
| GENERAL DESCRIPTION OF TECHNIQUE | 1 |
| SPECIFIC TECHNIQUE | 3 |
| Case 1: $NS_x > 0$ and $NL_y > 0$ | 3 |
| Case 2: $NS_y > 0$ and $NL_x > 0$ | 4 |
| Case 3: $NS_x > 0$ and $NL_x > 0$ | 4 |
| Case 4: $NS_y > 0$ and $NL_y > 0$ | 6 |
| SUBROUTINE PROBDIF | 6 |
| SUMMARY | 7 |
| APPENDIX A | 9 |



| | |
|----------------|---------|
| SEARCHED | INDEXED |
| SERIALIZED | FILED |
| OCT 1964 | |
| FBI - NEW YORK | |
| A-1 | |

SUBROUTINE PROBDIF

INTRODUCTION

Given n independent points x_1, x_2, \dots, x_n from the density $f_x(\cdot)$ and m independent points y_1, y_2, \dots, y_m from the density $f_y(\cdot)$, we wish to determine the probability that

$$\begin{aligned}f_x(Z) &= g(Z) \\f_y(Z) &= g(Z-\mu) .\end{aligned}$$

That is, we wish to determine the probability that one density is shifted by μ from the other density. Since we do not want to assume anything about the density $g(\cdot)$, we will employ a nonparametric procedure.

GENERAL DESCRIPTION OF TECHNIQUE

If the x 's and y 's come from the same density, there are $(m+n)!/n!m!$ possible orderings of the points; and each ordering is equally likely. If there are n' x 's smaller than any y and m' y 's larger than any x , the probability that the x 's and y 's come from a different density is equal to the probability of obtaining any sequence that starts with n' or more x 's and ends with m' or more y 's,

As an example of this procedure, let us assume that there are 3 x samples 1.0, 2.0, and 3.0 and 3 y samples 3.5, 4.5, and 5.5. Given these examples, there are 20 possible orderings for the samples and these orderings are listed in Table 1. The probability that the x points are μ less than the y points is given in Table 2. For $\mu=0$ the sequence has 3 x 's followed by 3 y 's. Only sequence #1 satisfies this condition and its probability of occurrence is 0.05. Thus, the probability that the samples do not come from the same density is one minus this probability or 0.95. For $\mu=1$, the sequence starts with 2 x 's and ends with 2 y 's. Sequences #1 and #2 satisfy this condition, their probability of occurrence is 0.10, and the probability that the samples do not come from the same density is 0.90. For $\mu=2$, the sequence starts with an x and ends with a y . Sequences #1, #2, #3, #5, #6, and #8 satisfy this condition, their probability is 0.3, and thus the probability that the samples do not come from the same density is 0.7. When the sequence starts with y 's and ends in x 's, the probability of occurrence of the sequences equals the probability that the samples do not come from the same density; i.e., one does not have to subtract the sequence probabilities from 1.

When the number of samples is a moderate number, it becomes impractical to enumerate all possible sequences. For instance, if $n=m=15$, the number of

Table 1 - Possible Sequences

| Sequence # | Sequence |
|------------|----------|
| 1 | xxxxyy |
| 2 | xxxyyy |
| 3 | xyyyxy |
| 4 | xyyyxx |
| 5 | xyxxyy |
| 6 | xyxyxy |
| 7 | xyxyxx |
| 8 | xyyyxy |
| 9 | xyyyxx |
| 10 | xyyyxx |
| 11 | yxxxxy |
| 12 | yxxxyx |
| 13 | yxxxyx |
| 14 | yxyxxy |
| 15 | yxyxyx |
| 16 | yxyyxx |
| 17 | yyxxxxy |
| 18 | yyxxyx |
| 19 | yyxyxx |
| 20 | yyyxxx |

Table 2 - Probability that the density of the x points shifted by μ is less than the density of the y points.

| μ | Probability |
|-------|-------------|
| 0 | .95 |
| 1 | .90 |
| 2 | .70 |
| 3 | .30 |
| 4 | .10 |
| 5 | .05 |

possible sequences is greater than 10^8 . Consequently, we will discuss a different technique for calculating the probability of the various sequences without enumerating them. This technique will also handle the cases where the smallest and largest samples come from the same set of points.

SPECIFIC TECHNIQUE

Let us first define NS_x as the number of x's which are smaller than all of the y's and NS_y as the number of y's which are smaller than all of the x's. Similarly, define NL_x as the number of x's which are larger than all of the y's and NL_y as the number of y's which are larger than all of the x's. Note that if $NS_x > 0$, then $NS_y = 0$; and if $NS_y > 0$, then $NS_x = 0$. Similarly, if $NL_x > 0$, then $NL_y = 0$; and if $NL_y > 0$, then $NL_x = 0$. Based on these four values (NS_x , NS_y , NL_x and NL_y), four different cases are possible:

1. $NS_x > 0$ and $NL_y > 0$,
2. $NS_y > 0$ and $NL_x > 0$,
3. $NS_x > 0$ and $NL_x > 0$,
4. $NS_y > 0$ and $NL_y > 0$.

Each of these cases will now be discussed and a formula will be generated to calculate the probability that the x data set comes from a density which has a smaller location parameter than the y density.

Case 1: $NS_x > 0$ and $NL_y > 0$

If $NS_x = n$, that is all the x's are less than all the y's, the probability of obtaining the first x in the sequence is $n/(m+n)$, the probability of obtaining the second x in the sequence is $(n-1)/(m+n-1)$, and the probability of the third x is $(n-2)/(m+n-2)$, and the probability of the i-th x is $(n+1-i)/(m+n+1-i)$. Consequently, the probability of this sequence is

$$\frac{n}{m+n} \cdot \frac{n-1}{m+n-1} \cdot \frac{n-2}{m+n-2} \cdots \frac{1}{m+1} = \frac{n!m!}{(m+n)!} \quad (1)$$

If $NS_x < n$, let i be in the range $NS_x \leq i < n$; and let i be the exact number of x's which start a sequence. The probability that exactly i x's start a sequence follows from the above argument and equals

$$\frac{n}{m+n} \cdot \frac{n-1}{m+n-1} \cdots \frac{n+1-i}{m+n+1-i} = \frac{n!}{(n-i)!} \frac{(m+n-i)!}{(m+n)!} \quad (2)$$

Since the sequence starts with exactly i x's, the next sample must be a y; and the probability of this y is

$$\frac{m}{(m+n-i)} \quad (3)$$

So far we have considered $i+1$ samples: i samples coming from the x set and 1 sample coming from the y set. Let us now assume that the sequence ends with exactly j y samples. This sequence of y samples must be preceded by an x, and the probability of this x is

$$\frac{n-i}{m+n-i-1} \quad (4)$$

Finally the probability of ending with j y samples is

$$\frac{m-1}{m+n-i-2} \cdot \frac{m-2}{m+n-i-3} \cdots \frac{m-j}{m+n-i-j-1} = \frac{(m-1)!}{(m-j-1)!} \frac{(m+n-i-j-2)!}{(m+n-i-2)!} \quad (5)$$

Thus, the probability P_{ij} of all the sequences that start with exactly i x 's and end with exactly j y 's is the product of expressions (2), (3), (4), and (5), i.e.,

$$P_{ij} = \frac{n!}{(n-i)!} \frac{(m+n-i)!}{(m+n)!} \frac{m}{(m+n-i)} \frac{n-i}{(m+n-i-1)} \frac{(m-1)!}{(m-j-1)!} \frac{(m+n-i-j-2)!}{(m+n-i-2)!} \quad (6)$$

Simplifying this expression, one obtains

$$P_{ij} = \frac{n!m!}{(m+n)!} \frac{1}{(n-i-1)!} \frac{(m+n-i-j-2)!}{(m-j-1)!} \quad (7)$$

For example, if $m=n=3$, $i=2$, and $j=1$, $P_{ij} = 1/20$, which corresponds to sequence #3 in Table 1. If $m=n=3$, $i=1$, and $j=1$; $P_{ij} = 2/20$, which corresponds to sequences #6 and #8 in Table 1. Equation (7) is valid for $NS_x \leq i < n$ and $NL_y \leq j < m$. Thus, the probability of obtaining sequences which start with at least NS_x x 's and end with at least NL_y y 's can be obtained by summing Eq. (7) over the proper ranges of i and j and adding expression (1) to this sum. Consequently, by subtracting this probability from 1, one obtains the probability that the x points come from a density which has a smaller location parameter than the y density has. The desired probability is

$$P = 1 - \frac{n!m!}{(m+n)!} \left\{ 1 + \sum_{i=NS_x}^{n-1} \frac{1}{(n-i-1)!} \left[\sum_{j=NL_y}^{m-1} \frac{(m+n-i-j-2)!}{(m-j-1)!} \right] \right\} \quad (8)$$

Case 2: $NS_y > 0$ and $NL_x > 0$

This case is the dual of case 1. If we let $NS_y \leq i < m$ and $NL_x \leq j < n$, the revised Eq. (8) would yield the probability that the y 's are less than the x 's. The probability that the x 's are less than the y 's is 1 minus this probability or

$$P = \frac{n!m!}{(m+n)!} \left\{ 1 + \sum_{i=NS_y}^{m-1} \frac{1}{(m-i-1)!} \left[\sum_{j=NL_x}^{n-1} \frac{(m+n-i-j-2)!}{(n-j-1)!} \right] \right\} \quad (9)$$

Case 3: $NS_x > 0$ and $NL_x > 0$

This case is likely to occur if $n \gg m$. If $NS_x = NL_x$, the desired probability is

$$P = 0.5 \quad .$$

(10)

If $NS_x \neq NL_x$, let $NS = \text{Minimum}(NS_x, NL_x)$ and let $NL = \text{Maximum}(NS_x, NL_x)$. Then, the sequences of interest are those sequences which start with NL or more x's and end with either NS or less x's or any number of y's. The probability that a sequence starts with n x's and ends with m y's is given by expression (1). Let i be in the range $NL \leq i < n$ and let i be the exact number of x's which start a sequence. The probability that exactly i x's start a sequence is given by (2). Since the sequence starts with exactly i x's, the next sample must be a y; and the probability of this y is given by (3). So far we have considered $i+1$ samples: i samples coming from the x set and 1 sample coming from the y set. Let us now assume that the sequence ends with exactly j x samples. This sequence of x samples must be preceded by a y, and the probability of this y sample is

$$\frac{m-1}{m+n-i-1} \quad . \quad (11)$$

The probability of ending with j x samples is

$$\frac{n-i}{m+n-i-2} \cdot \frac{n-i-1}{m+n-i-3} \dots \frac{n-i-j+1}{m+n-i-j-1} = \frac{(n-i)!}{(n-i-j)!} \frac{(m+n-i-j-2)!}{(m+n-i-2)!} \quad . \quad (12)$$

The possible values of j are from 1 to N_1 where N_1 is the minimum of NS and $(n-i)$. The probability of starting a sequence with exactly i x's and ending it with exactly j x's is the product of (2), (3), (11), and (12)

$$\frac{n!}{(n-i)!} \frac{(m+n-i)!}{(m+n)!} \frac{m}{(m+n-i)} \frac{m-1}{(m+n-i-1)} \frac{(n-i)!}{(n-i-j)!} \frac{(m+n-i-j-2)!}{(m+n-i-2)!} \quad (13)$$

which simplifies to

$$\frac{n!m(m-1)}{(m+n)!} \frac{(m+n-i-j-2)!}{(n-i-j)!} \quad . \quad (14)$$

The sequences of interest which start with exactly i x's can also end with j y samples and this probability is given by (7). Combining (7) and (14) and summing over the proper values of i and j , one obtains

$$P' = \frac{n!m!}{(m+n)!} + \sum_{i=N_L}^{n-1} \left\{ \frac{n!m(m-1)}{(m+n)!} \sum_{j=1}^{N_1} \frac{(m+n-i-j-2)!}{(n-i-j)!} \right. \\ \left. + \frac{n!m!}{(m+n)!} \frac{1}{(n-i-1)!} \sum_{j=1}^{m-1} \frac{(m+n-i-j-2)!}{(m-j-i)!} \right\} \quad . \quad (15)$$

Rearranging terms, one obtains

$$P' = \frac{m!n!}{(m+n)!} \left\{ 1 + \sum_{i=NL}^{n-1} \left[\frac{1}{(m-2)!} \sum_{j=1}^{N_1} \frac{(m+n-i-j-2)!}{(n-i-j)!} + \frac{1}{(n-i-1)!} \sum_{j=1}^{m-1} \frac{(m+n-i-j-2)!}{(m-j-1)!} \right] \right\} . \quad (16)$$

Finally, if $NS_x > NL_x$

$$P = 1 - P' ; \quad (17)$$

and if $NL_x > NS_x$

$$P = P' . \quad (18)$$

Case 4: $NS_y > 0$ and $NL_y > 0$

This case is the dual of case 3. Letting $NS = \text{Minimum}(NS_y, NL_y)$, $NL = \text{Maximum}(NS_y, NL_y)$, and N_2 be the minimum of NS and $(m-1)$, the probability P' is given by

$$P' = \frac{m!n!}{(m+n)!} \left\{ 1 + \sum_{i=NL}^{m-1} \left[\frac{1}{(n-2)!} \sum_{j=1}^{N_2} \frac{(m+n-i-j-2)!}{(m-i-j)!} + \frac{1}{(m-i-1)!} \sum_{j=1}^{n-1} \frac{(m+n-i-j-2)!}{(n-j-1)!} \right] \right\} . \quad (19)$$

If $NS_y > NL_y$,

$$P = P' ; \quad (20)$$

and if $NL_y > NS_y$

$$P = 1 - P' . \quad (21)$$

SUBROUTINE PROBDIF

Depending on the values of NS_x , NS_y , NL_x , and NL_y , the desired probability is given in Eq. (8), Eq. (9), Eqs. (16-18), or Eqs. (19-21). All of these equations involve double summations over factorials and the computation time will vary as n^3 (or m^3) if these formula are implemented

directly. However, since the terms involve factorials, iterative formulas can be developed and the computation time can be made proportional to n^2 . Such a technique was incorporated into subroutine PROBDIF which calculates the desired probability.

The calling sequence for the subroutine is

```
CALL PROBDIF (nx,x,ny,y,xdb,prob,iset)
```

where

nx is number of points from the x distribution
x(.) are points from the x distribution
ny is number of points from the y distribution
y(.) are points from the y distribution
xdb is the db difference to be tested
prob is the probability of a xdb difference
iset=1 indicates that points have already been
ordered from smallest to largest.

Note, if successive calls are made to PROBDIF with different values of xdb, iset should be set to 1 after the first call to minimize computation time. A Fortran listing of PROBDIF is given in appendix A.

SUMMARY

Given two sets of data, this report describes a technique for determining the probability that the first density is a shifted version of the second density. We assumed nothing about the density and used a nonparametric procedure based on the smallest and largest samples of the combined data set. A computer program was written to calculate the desired probability.

APPENDIX A

```

subroutine probdif(nx,x.ny,y,xdb,prob,iset)
c
c   calculates probability that x distribution is x dB less
c   than y distribution
c
c   nx is number of points from x distribution
c   x(.) are points from x distribution
c   ny is number of points from y distribution
c   y(.) are points from y distribution
c   xdb is the db difference to be tested
c   prob is the probability of a xdb difference
c   iset=1 indicates that points have already been ordered
c
dimension x(100),y(100),xo(100),yo(100)
if (iset.eq.1) go to 70
n=nx+ny
c
c   ordering of x points
c
do 5 i=1,nx
5   xo(i)=x(i)

if (nx.eq.1) go to 30

do 20 i=1,nx-1
do 10 j=i+1,nx
if (xo(i).lt.xo(j)) go to 10
t=xo(j)
xo(j)=xo(i)
xo(i)=t
10 continue
20 continue
c
c   ordering of the y points
c
do 35 i=1,ny
35   yo(i)=y(i)

if (ny.eq.1) go to 70

```

```

do 60 i=1,ny-1
do 40 j=i+1,ny
if (yo(i).lt.yo(j)) go to 40
t=yo(j)
yo(j)=yo(i)
yo(i)=t
40 continue
60 continue

70 continue
c
c      nsmallx is the # of x points less than yo(1)-xdb
c      ns малы is the # of y points less than xo(1)+xdb
c
nsmallx=0
ns малы=0
pl=1.0

do 80 i=1,nx
if (xo(i).gt.yo(1)-xdb) go to 85
nsmallx=nsmallx+1
80 continue

85 if (nsmallx.eq.0) go to 95
go to 120

95 do 100 j=1,ny
if (xo(1).lt.yo(j)-xdb) go to 105
ns малы=ns малы+1
100 continue

105 continue
c
c      nlargey is the # of y points greater than xo(nx)-xdb
c      nlargex is the # of x points greater than yo(ny)-xdb
c
120 nlargey=0
nlargex=0

do 130 j=1,ny
jj=ny+1-j
if (yo(jj).lt.xo(nx)-xdb) go to 135
nlargey=nlargey+1
130 continue

135 if (nlargey.eq.0) go to 145
go to 170

```

```

145 do 150 i=1,nx
    ii=nx+1-i
    if (yo(ny).gt.xo(ii)+xdb) go to 155
    nlargex=nlargex+1
150 continue

155 continue

170 continue

c
c      calculation of probabillity
c
    prob=0.0
    nxp=nx+1
    nyp=ny+1
    np=n+1
    if (nsmallx.gt.0.and.nlargey.gt.0) go to 200
    if (nsmally.gt.0.and.nlargex.gt.0) go to 300
    if (nsmallx.gt.0.and.nlargex.gt.0) go to 400
    if (nsmally.gt.0.and.nlargey.gt.0) go to 500
    print 50
50   format ('          ***** impossible condition *****')
    return

c
c      case #1  nsmallx>0 and nlargey>0
c
200  f=1.

    do 250 i=nsmallx,nx
    if (i.gt.nsmallx) go to 210

    do 205 j=1,nsmallx
    nxp=nxp-1
    np=np-1
    f=(f*nxp)/np
205  continue

    if (i.lt.nx) go to 210
208  h=1.
    go to 245
210  if (i.eq.nx) go to 208
    nxp=nxp-1
    np=np-1
    f=(f*nxp)/np
    m=np-1
    nyp=nyp
    g=(1.*nyp)/m

    do 240 j=nlargey,ny-1
    if (j.gt.nlargey) go to 230

```

```

do 225 k=1,nlargey
nyp=nyp-1
m=m-1
g=(g*nyp)/m
h=g
225 continue

go to 240
230 nyp=nyp-1
m=m-1
g=(g*nyp)/m
h=h+g
240 continue

245 prob=prob+f*h
250 continue

prob=1.-prob
return

o
c case #2 nsmally>0 and nlargex>0
o
300 continue
f=1.

do 350 i=nsmally.ny
if (i.gt.nsmally) go to 310

do 305 j=1,nsmally
nyp=nyp-1
np=np-1
f=(f*nyp)/np
305 continue

if (i.lt.ny) go to 310
308 h=1.
go to 345
310 if (i.eq.ny) go to 308
nyp=nyp-1
np=np-1
f=(f*nyp)/np
320 m=np-1
nyp=nyp-1
g=(1.*nyp)/m

do 340 j=nlargex,nx-1
if (j.gt.nlargex) go to 330

```

```

do 325 k=1,nlargex
nxp=nxp-1
m=m-1
g=(g*nxp)/m
h=g
325 continue

go to 340
330 nxp=nxp-1
m=m-1
g=(g*nxp)/m
h=h+g
340 continue

345 prob=prob+f*h
350 continue

return

c
c case #3 nsmallx>0 and nlargey>0
c
400 continue
if (nsmallx.ne.nlargex) go to 401
prob=0.5
return
401 ns=min(nsmallx,nlargex)
nl=max(nsmallx,nlargex)
f=1.0

do 450 i=nl,nx
if (i.gt.nl) go to 410

do 405 j=1,nl
nxp=nxp-1
np=np-1
f=(f*nxp)/np
405 continue

if (i.lt.nx) go to 420
408 h=1.
go to 445
410 nxp=nxp-1
np=np-1
f=(f*nxp)/np
if (i.eq.nx) go to 408
420 h=0.
m=np-1
nyp=ny
g=(1.*nyp)/m
if (nyp.eq.1) go to 422
m=m-1
nyp=nyp-1
g=(g*nyp)/m

```

```

422  f1=1.0
      nxpp=nxp

      do 425 j=1,ns
        nxpp=nxpp-1
        if (nxpp.eq.0) go to 426
        m=m-1
        f1=(f1*nxpp)/m
        h=h+g*f1
425  continue

426  m=np-1
      g=(nxp-1.)/m
      nyp=ny
      m=m-1
      g=(g*nyp)/m
      f1=1.

      do 440 j=1,ny-1
        nyp=nyp-1
        m=m-1
        f1=(f1*nyp)/m
        h=h+g*f1
440  continue

445  prob=prob+f*h
450  continue

      if (nsmallx.gt.nlargex) prob=1.0-prob
      return
o
o      case #4 nsmally>0 and nlargey>0
o
500  continue
      if (nsmally.ne.nlargey) go to 501
      prob=0.5
      return
501  ns=min(nsmally,nlargey)
      nl=max(nsmally,nlargey)
      f=1.0

      do 550 i=nl,ny
        if (i.gt.nl) go to 510

        do 505 j=1,nl
          nyp=nyp-1
          np=np-1
          f=(f*nyp)/np
505  continue

```



```

      if (i.lt.ny) go to 520
508   h=1.
      go to 545
510   nyp=nyp-1
      np=np-1
      f=(f*nyp)/np
      if (i.eq.ny) go to 508
520   h=0.
      m=np-1
      nxp=nx
      g=(1.*nxp)/m
      if (nxp.eq.1) go to 522
      m=m-1
      nxp=nxp-1
      g=(g*nxp)/m
522   fl=1.0
      nypp=nyp

      do 525 j=1,ns
      nypp=nypp-1
      if (nypp.eq.0) go to 526
      m=m-1
      fl=(fl*nypp)/m
      h=h+g*fl
525   continue

526   m=np-1
      g=(nyp-1.)/m
      nxp=nx
      m=m-1
      g=(g*nxp)/m
      fl=1.

      do 540 j=1,nx-1
      nxp=nxp-1
      m=m-1
      fl=(fl*nxp)/m
      h=h+g*fl
540   continue

545   prob=prob+f*h
550   continue
      if (nsmally.lt.nlargey) prob=1.0-prob
      return
      end

```